

# Package ‘boxfilter’

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**Type** Package

**Title** Filter Noisy Data

**Version** 0.1

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**Description** Noise filter based on determining the proportion of neighboring points. A false point will be rejected if it has only few neighbors, but accepted if the proportion of neighbors in a rectangular frame is high. The size of the rectangular frame as well as the cut-off value, i.e. of a minimum proportion of neighbor points, may be supplied or can be calculated automatically. Originally designed for the cleaning of heart rates, but suitable for filtering any slowly-changing physiological variable. For more information see Signer (2010)<[doi:10.1111/j.2041-210X.2009.00010.x](https://doi.org/10.1111/j.2041-210X.2009.00010.x)>.

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** true

**Imports** ggplot2, gridExtra

**Depends** R (>= 3.50)

**Suggests** knitr, rmarkdown

**VignetteBuilder** knitr

**NeedsCompilation** no

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**Repository** CRAN

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<b>boxclip</b>	<i>Boxfilter Main Function</i>
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**Description**

Filters noise from data (e.g. heartrates) using x for x-axis data and y for y-axis, based on the proportion of neighbors of each point in a surrounding box of width and height (these may be determined automatically). It discards all data points that have less than a proportion of clipit neighbors.

**Usage**

```
boxclip(x, y, clipit=NULL, QI=NULL, width=NULL, height=NULL,
        miny=10, plotit=TRUE, histo=FALSE)
```

**Arguments**

x	The x-axis of data, a datetime for example. Optional. If x=NULL x<-1:length(y) will be generated.
y	The y-axis of data, probably noisy. Required.
clipit	Optional. Y-values with less than a proportion of clipit neighbors will be discarded. If clipit is omitted it is set equal to the first trough in the neighbor proportion histogram. ci
QI	Optional. An integer quality index for each data point.
width	Optional. The width of the box. If width is omitted it will be generated from floor(length(x)*0.01).
height	Optional. The height of the box. If height is omitted it will be generated from floor(mean(y,na.rm=T)/4).
miny	The minimum y-value expected. Defaults to 10. Anything below miny is discarded.
plotit	Optional. If TRUE show a graph of the original and filtered data.
histo	Optional. If TRUE also show a histogram of the neighboring points.

**Details**

Boxfilter mimics the human criterion of self-similarity. Data points with many neighbors are more trustworthy.

**Value**

x	Original x-axis data
y	Original y-axis data
filtered	Filtered data. Discarded data points are set to NA.
neighbors	Proportion of neighbors of each point.

**Note**

To store only filtered data, use e.g.:

```
bc=boxclip(x,y) data=data.frame(x=bc$x, hrf=bc$filtered) data=na.omit(data) write.csv(data,file="myheartrates.csv")
```

**Author(s)**

Thomas Ruf (thomas.p.ruf@me.com)

**See Also**

clipview

**Examples**

```
data("wb_month.RData")
data ("ibex_hr.RData")

x=wb_month$x
y=wb_month$hr

myclip=boxclip(x,y,clipit=0.25,width=15, histo=TRUE)
summary(myclip)

r=seq(1,28400,by=4)
myclip=boxclip(ibex_hr$Time[r],ibex_hr$Heartrate[r],0.45)
summary(myclip)
store(myclip)
```

---

clipview

*Get View Of Different clipit Values*

---

**Description**

Gives a view of the original data, a histogram of the neighbors, and results of four values of clipit. The cutoff-values clipit are determined by a sequence of length .

**Usage**

```
clipview(x, y, clipit = NULL, width = NULL, height = NULL, miny=10)
```

## Arguments

x	The x-axis of data, a datetime for example. Optional. If x=NULL x<-1:length(y) will be generated.
y	The y-axis of data, probably noisy. Required.
clipit	This function requires a sequence of four items <1, e.g. clipit=seq(0.1, 0.4, by=0.1)
width	Optional. The width of the box. If width is omitted it will be generated from floor(length(x)*0.01).
height	Optional. The height of the box. If height is omitted it will be generated from floor(mean(y,na.rm=T)/4).
miny	The minimum y-value expected. Defaults to 10. Anything below miny is discarded.

## Details

Note that a sequence is required for clipit here, while a scalar is required in boxclip().

## Value

There is no return value. Six graphs are generated, original, histogram, and four for each value of clipit.

## Author(s)

Thomas Ruf (thomas.p.ruf@me.com)

## See Also

boxclip()

## Examples

```
data("wb_year.RData")

r=seq(1,54179,by=5)
x=wb_year$x[r]
y=wb_year$y[r]

clipview(x,y,clipit=seq(0.2,0.5,0.1))
```

---

ibex\_hr

*One year of heart rates of a capricorn free-living in the alps.*

---

### Description

Heart rates were obtained using acoustic loggers in the rumen.

### Usage

```
data("ibex_hr")
```

### Format

A data frame with 28454 observations on the following 2 variables.

Time datetime

Heartrate in bpm

### References

Signer, C., Ruf, T., & Arnold, W. (2011). Hypometabolism and basking: The strategies of Alpine ibex to endure harsh over-wintering conditions. *Functional Ecology*, 25(3), 537-547.

### Examples

```
data(ibex_hr)
```

---

store

*Store data*

---

### Description

Stores original and filtered data.

### Usage

```
store(object)
```

### Arguments

object            Must be of class "boxclip", resulting from boxclip.

### Value

returns nothing

**Author(s)**

Thomas Ruf (thomas.p.ruf@me.com)

**Examples**

```
data ("wb_month.RData")
x=wb_month$x
y=wb_month$hr

myclip=boxclip(x,y,clipit=0.25,width=15)
summary(myclip)
store(myclip)
```

**summary.boxclip**      *Summarize boxclip results*

**Description**

Summary method for class `boxclip`.

**Usage**

```
## S3 method for class 'boxclip'
summary(object,...)
```

**Arguments**

<code>object</code>	an object of class <code>boxclip</code>
<code>...</code>	currently, no other arguments are required.

**Value**

`summary.boxclip` prints the following items:

<code>clipit</code>	Cut-off value. Only points with a proportion $\geq$ <code>clipit</code> will be retained.
<code>width</code>	Rectangle width in x-units. Filter criterion is the proportion of data points inside the rectangle.
<code>height</code>	Rectangle height in y-units. Filter criterion is the proportion of data points inside the rectangle.
<code>full</code>	Number of original data.
<code>remaining</code>	Number of data remaining.
<code>rest</code>	Percentage remaining.

**Author(s)**

Thomas Ruf <thomas.p.ruf@me.com>

**References**

Signer (2010) <doi: 10.1111/j.2041-210X.2009.00010.x>

**See Also**

[boxclip](#)

**Examples**

```
data ("wb_month.RData")  
  
x=wb_month$x  
y=wb_month$hr  
  
myclip=boxclip(x,y,clipit=0.25,width=15)  
summary(myclip)
```

---

wb\_month

*One month of heart rates and their quality in a wild boar. Quality was assessed by Star-Oddi, Iceland.*

---

**Description**

Heart rates were obtained from DST centi- HRT, Star-Oddi, Gardabaer, Iceland.

**Usage**

```
data("wb_month")
```

**Format**

A data frame with 3720 observations on the following 3 variables.

- x x-axis, datetime
- hr y-axis, heart rate
- QI quality index (0-3) of the signal

**Examples**

```
data(wb_month)
```

---

*wb\_year*                    *One year of heart rates of a wild boar female.*

---

### Description

Heart rates were obtained from DST centi- HRT, Star-Oddi, Gardabaer, Iceland.

### Usage

```
data("wb_year")
```

### Format

A data frame with 3720 observations on the following 3 variables.

- x x-axis, datetime
- y y-axis, heart rate

### References

Ruf, T., Vetter, S. G., Painer, J., Stalder, G., & Bieber, C. (2021). Atypical for northern ungulates, energy metabolism is lowest during summer in female wild boars (*Sus scrofa*). *Scientific Reports*, 11(1), 1-12.

### Examples

```
data(wb_year)
```

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